

SCORE Search Results Details for Application 10063597 and Search Result us-10-063-597- 90.rspt.

[Score Home Page](#) [Retrieve Application List](#) [SCORE System Overview](#) [SCORE FAQ](#) [Comments / Suggestions](#)

This page gives you Search Results detail for the Application 10063597 and Search Result us-10-063-597-90.rspt.

[start](#)

[Go Back to previous page](#)

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: June 22, 2004, 09:53:57 ; Search time 39 Seconds
(without alignments)
1132.629 Million cell updates/sec

Title: US-10-063-597-90
Perfect score: 716
Sequence: 1 MGRVSGLVPSRPLTLLAHLV.....PAVTEMALFVTVFGLKKPF 140

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1017041 seqs, 315518202 residues

Total number of hits satisfying chosen parameters: 1017041

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : SPTREMBL_25:
1: sp_archea:
2: sp_bacteria:
3: sp_fungi:
4: sp_human:
5: sp_invertebrate:
6: sp_mammal:
7: sp_mhc:
8: sp_organelle:
9: sp_phage:
10: sp_plant:
11: sp_rat:
12: sp_virus:
13: sp_vertebrate:
14: sp_unclassified:
15: sp_rvirus:
16: sp_bacteriap:
17: sp_archeap:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Score	Match	Length	DB	ID	Description
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EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	689	SEQ ID NO:90 PRO1268 95% 99%	US-PGPUB; USPAT; EPO; DERWENT	AND	OFF	2006/06/25 18:19
L2	915	EATON GODDARD WOOD WATANABE	US-PGPUB; USPAT; EPO; DERWENT	AND	OFF	2006/06/25 18:20